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APPENDIX TO AMENDMENT OF MARCH 6, 2003

Version with Markings to Show Changes Made

Amendments to the Specification

The paragraph on page 47, beginning on line 5:

—A sample of [the] human MI [preparation described (about 6.5 µg)] was subjected to amino-terminal amino acid sequencing through 18 cycles, using the methods described in Example 2. The initial yield was 135 pmol and the repetitive yield was 94%. The major sequence obtained exactly matched that predicted from the nucleotide sequence for the mature human MI gene (Example 3; Figure 2).—

Amendments to the Claims

40. [A] An isolated DNA encoding a polypeptide product consisting of an amino acid sequence selected (according to the numbering as presented in Figure 2) from the group consisting of:

- (a) amino acids -26 through 194;
- (b) amino acids 1 through 194; and
- (c) [a polypeptide of subpart (a), lacking residues -26 through -1 and having a methionyl residue at position -1; and]

[(d)] a polypeptide of subpart (b), having [a] an additional methionyl residue at position -1.

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